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New Atty. Docket No: 37629-0004
Old Atty. Docket No. 047744/0106

#14/D

*In re patent application of
KNAPPIK, Achim et al.*

Application No.: 09/490,324

Group Art Unit: 1631

Filed: January 24, 2000

Examiner: M. Zeman

For: PROTEIN/ (POLY) PEPTIDE LIBRARIES

NE

AMENDMENT

Commissioner for Patents
Washington, D.C. 20231

*Duplicate
of #11
except for
seq. listing*

Sir:

In response to the Office Action mailed May 7, 2002, Applicants respectfully request that the following amendments be entered into the application:

IN THE SPECIFICATION:

Page 18, line 4, paragraph 2: Please amend as follows:

Figs. 2A-2G: Alignment of consensus sequences designed for each subgroup (amino acid residues are shown with their standard one-letter abbreviation). (2A-2B) (SEQ ID NOS 28-31, respectively) kappa sequences, (2C-2D) (SEQ ID NOS 32-34, respectively) lambda sequences and (2E-2G) (SEQ ID NOS 35-41, respectively), heavy chain sequences. The positions are numbered according to Kabat (1991). In order to maximize homology in the alignment, gaps (-) have been introduced in the sequence at certain positions.

Page 18, line 10, paragraph 3: Please amend as follows:

Figs. 3A-3K: Gene sequences (SEQ ID NOS 42, 44, 46 and 48, respectively) of the synthetic V kappa consensus genes. The corresponding amino acid sequences (SEQ ID NOS 43, 45, 47 and 49, respectively) (see Figs. 2A-2B) as well as the unique cleavage sites are also shown.